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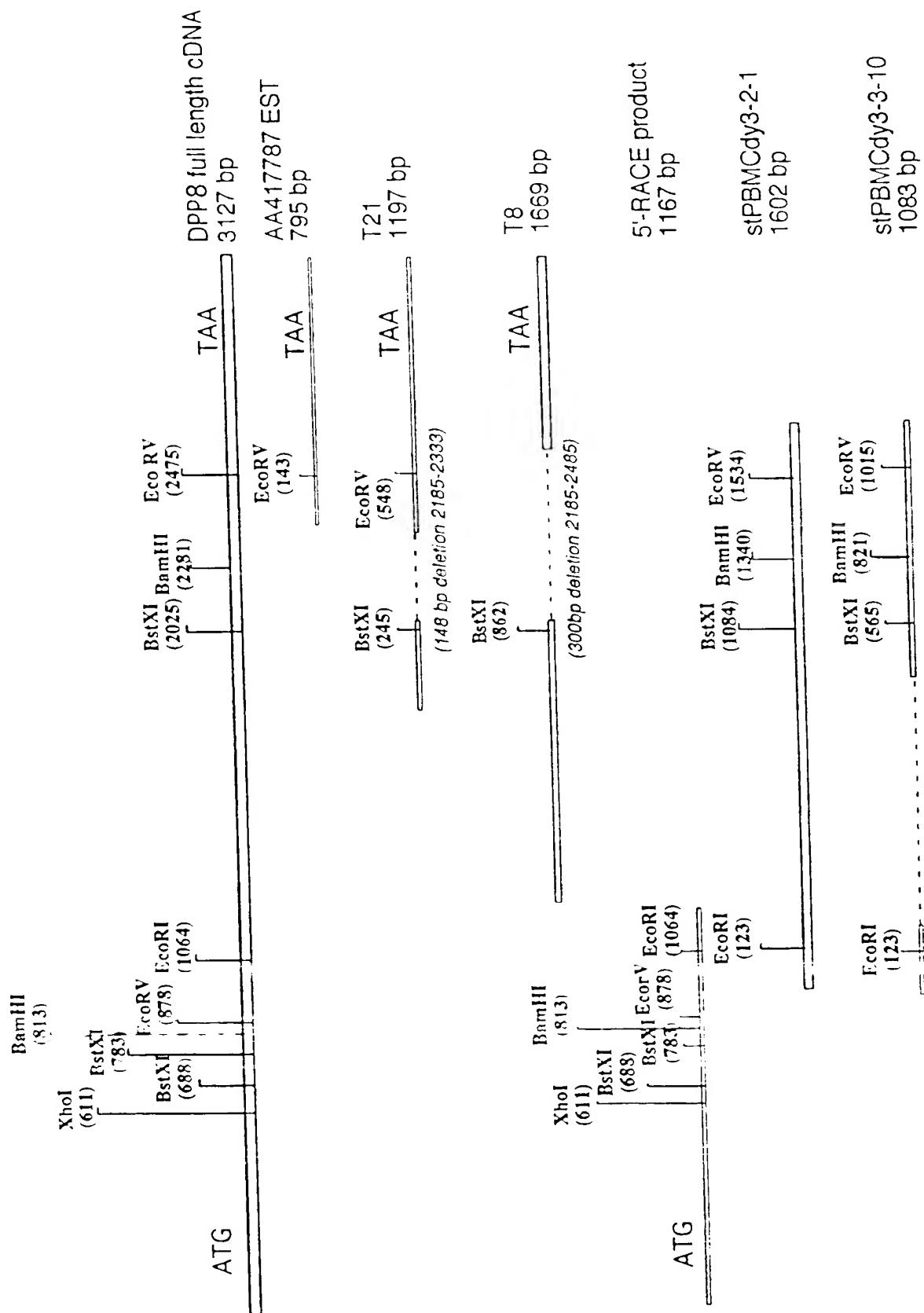


FIGURE 1

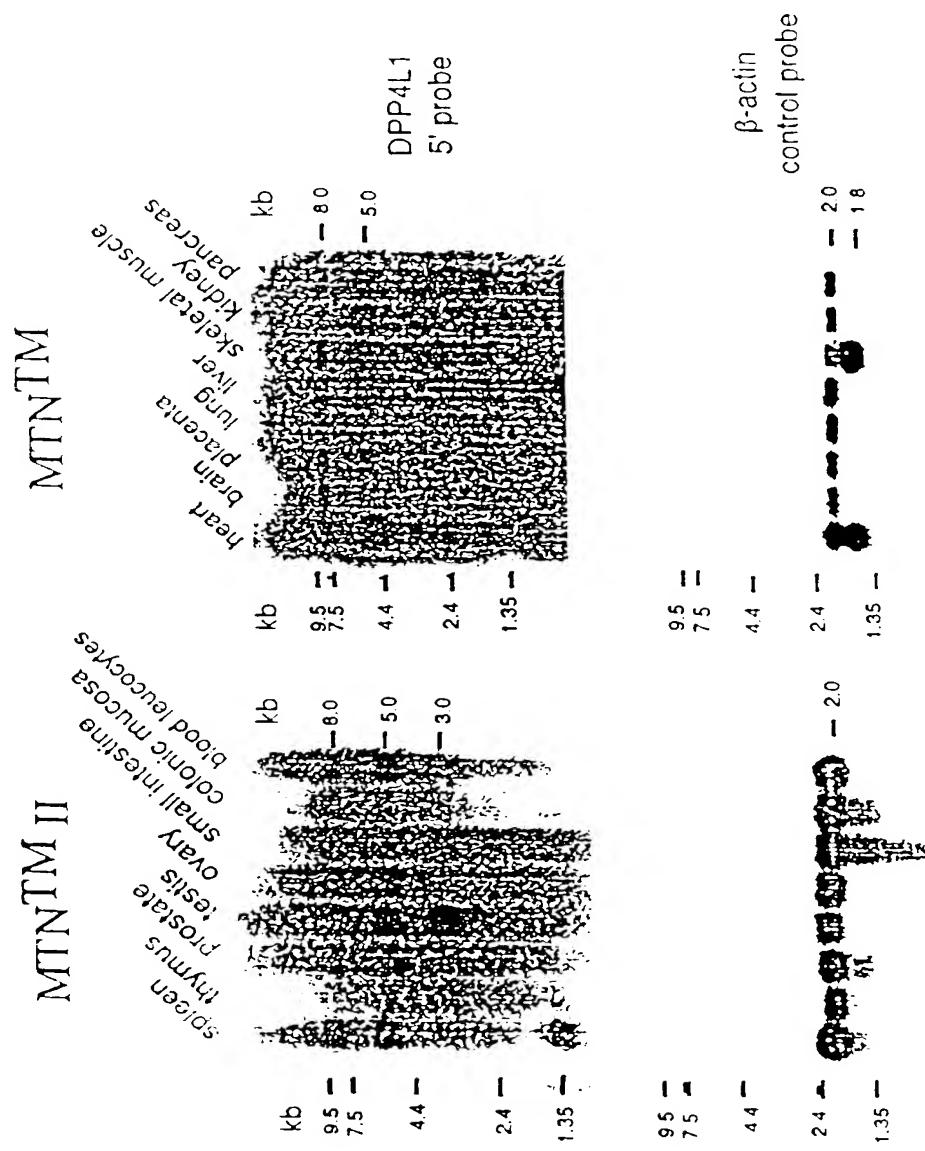
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 101 AGCCACTGCACCAAGGACCCGACTGAGCGCGGCGACGCTGAAAGCAGCGCTGAGGATATTGAAACTGCGACTGTGAGGAGAATTTGAATCACAGGA 200
 201 AGGAAAATGCAACATGGCAGGACGAACTGAAACAGAACAGCTGGGTTGAGATATTGAAACTGCGACTGTGAGGAGAATTTGAATCACAGGA 300
 K A A A K E T E O L G V E I F E T A D C E E N I E S Q D R 29
 301 CCTAAATGGGACCTTTTATGTTGAGGGTATTCCTGGGTCAGCTTAAAGCTGCTTGGCGATAACAGAAMATACTACGCTACATCATGGCTAAGG 400
 30 P K L E P F Y V E R Y S W S O L K K L A D T R K Y H C Y H M A K A 63
 401 CACACATGATTCTAGTTCTGAAAGAGGAATGATCCAGATGGACCTCATCACAGAACAGATCTATTACCTGGCAATGCTGGTGAGAACAGAGAAA 500
 63 P H D F H F V K R N D P D G P H S D R I Y Y L A M S G E N R E N T 96
 501 ACTCTTTATTCGAAATTCGAAACATCAATAGAGGACGACTTAAATGCTCTTGGCTTGGATCTTCAAGGAACTGGACTAT 600
 96 L F Y S E I P K T I N R A A V L M L S W K P L L D L F Q A T L D Y 129
 601 CGAAATGATTCGAGAGAGAACTATTAGAGAAAGAAAACCGATGGAAAGCTGGAAATTGCTCTTCAAGGAACTGGAAACAGGAACTGGACTAT 700
 130 G M Y S R E E E L L R E R K R I G T V G I A S Y D Y H Q G S G T F L 163
 701 TGTTCAAGGGCTAGTGGAAATTATCACGTAAGGATGGGGCCACAAAGGATTAGCGAACACCTTAAGGCCAACTGAAACTAGTGGAAACTAGTTG 800
 163 F O A G S G I Y H V K D G G G P Q G F T Q Q P R N L V E T S C P 196
 801 CAACATACGGATGGATCCAAATTATGCCCGGCTGATCAGACTGGATCTGCTTACATAGCAACGATATTGGATCTAACATGTAACAGAGAA 900
 196 N I R D P K L C P A D P D W I A F I H S N D I W I S N I V T R E 229
 901 GAAGGAGACTCACTTATGCAACATGAGCTAGCCACATGGAAAGAGATGCCAGATAGCTGGACTCTTCTCAAGAACAGATACTGATA 1000
 230 E R R L T Y V H N E L A N H E D A R S A G V A T F V L Q E E F D R 263
 1001 GATATTCTGGTATGGTGGTGGCAAAGCTGAAACAACTCCAGTGGTGGTAAATTCTTAAAGGATATGCTATATGAGAAATATGATGAACTGGTGA 1100
 263 Y S G Y W H C P K A E T T P S G G K I L R I L Y E E N D E S E V E 296
 1101 ATTATTCATTTACATCCCCATGTTGAAACAAAGGAGGGCAGATTCTTCCATTTCTAAACAGGTACAGCAATTCTAAAGTCACCTTAAAGATC 1200
 296 I I H V T S P K L E T R R A D S F R Y P K T G T A N P K V T F K H 329
 1201 TCAGAAAATATGATTGATGCTGAAAGGAGATCATAGATGTCATAGATANGGAACTAATTCAACCTTGGAGATCTTAAAGGAGATATATTG 1300
 330 S E I M Y D A E G R I I D V I D K E L T I Q P F E I L F E G V E Y I A 363
 1301 CCAGACCTGGATGACTCTGGGGAAAATATGCTTGGTCACTAGATAGCTGGCTCCAGACTGGCTACAGATAGTGGTGTCTACCTGAAATTATT 1400
 363 P A G H T P E G K Y A W S T I L L D R S O T R L Q I V L S I P E L F 396
 1401 TATCCCTAGAAGATGATGTTATGCAAGGAGACTATTGAGCTAGTGGCTGATTCGACGGCCACTAATTATCTATGAGAAACACAGACATC 1500
 396 I P V E D D V H E R Q R L I E S V P D S V T P L I F E E T T D I 429
 1501 TGGATAAAATACATGACATCTTACATGTTTCCCAGGAAAGTCAAGGAGGAAATTGAGTTTATTTTCCCTCTGAACTGAAACAGTTGGCTATT 1600
 430 W I N I H D I F H V F P Q S H E E E I E F I F A S E C K T G F R H L 463
 1601 TATACAAAATTACATCTATTAAAGGAAAGCAAATAAAACGATCCAGTGGTGGCTGCTGCTCAAGTGTGTTCAAGTGTCTATCAAGAGGAGAT 1700
 463 Y K I T S I L K E S K Y K R S S G G L P A P S D F K C P I K E E I 496
 1701 AGCAATTACAGTGGTGAATGGGAAGTCTGGCCGGCATGGATCTAAATCTCAAGTGTGATGAGCTAGAGGCTGGTATTTGAGGCAAAAGAC 1800
 496 A I T S G E W E V L G R H G S N I Q V D E V R R L V Y F E G T K D 529
 1801 TCCCTTCTAGAGCATCACCTGAGCTGAGCTGAGCTTACGTAATCTGGAGAGCTGACAGGCTGACTGACCTGGCTACTCACATTCTGGCATGCT 1900
 530 S P L E H H L Y V V S Y V N P G E V T R L T D R G Y S H S C C I S O 563
 1901 AGCACTGCTGACTCTTAAAGTAAGTATGAACTCACAAGAACTGGCTCTTACAGCTGAGCTGAGCTGAGCTGAGCTGAGCTGAGCTGAGCTG 2000
 563 H C D F F I S K Y S N Q K N P H C V S L Y K L L S S P D D P T C K 596
 2001 AACAAAGGAAATTGGGCCACCATTTGGATTCAAGGAGGCTCTCTCTGACTATACCTCTGAGGAAATTCTCTCTGAGGACTACTOGATTTACA 2100
 596 T K E F W A T I L D S A G P L P D Y T P P E I S F S E S T T G F T 629
 2101 TTCTATGGGATCTTACAGCTCATGATCTACACCTGAGGAAATTACCTCTGCTGCTGCTATATGGTGGTCTCAGTGGCTGGTGAATA 2200
 630 L Y G H L Y K P H D L Q P G K K Y P T V L F I Y G G P O V O L V N N 663
 2201 ATCGGTTAAAGGACTCAAGTATTCGGCTGAAATACCCCTACGGTCTCTAGGTATCTGCTGCTGAGCTGAGACACAGGGATCTCTACCGAGGGCT 2300
 663 R F K G V K Y F R L N T L A S L G Y V V V V I D N R G S C H R G L 696
 2301 TAAATTGAGGGCTTAAATATAAAATGGCTAAATAGAAATTGAGGATCAAGTGGAGACTCTAACATCTAGCTTCTGAGTATGATCTGAC 2400
 696 F F E G A P F Y E R G Q Z E I E D C V E G C L U Y L A S F T D P I D 729
 2401 TTAGATGCTGGCATCCACGGCTCTGAGGATACCTCTCCCTGAGGATACCTCTGGATGAGCTTACGAGGCTGAGCTGAGCTGAGCTGAGCTG 2500
 730 I D R V G T H [REDACTED] G Y L S L H A M L H O R S D I T R V A I A G A 763
 2501 CCCACTGACTCTGGATCTCTGATACAGGATACACGGAGCTGTTATGGTCTCTGAGGATCTCTGGATGAGCTGAGCTGAGCTGAGCTGAGCTG 2600
 763 P V T L W I F Y D T G Y T E R Y H G H P D O N E O G Y Y L G S V A 796
 2601 CATGCAAGGAGAAAGTCTCCCTGAAACAAATGTTACTGCTCTACATGCTGCTGAGGATGAGATGCTGAGCTGAGCTGAGCTGAGCTGAGCTG 2700
 796 M Q A E K F P S E P K R L L L H G P L E N V H F A R T S I L L 829
 2701 AGTTTTTACTGAGGGCTGAGGAAAGGCTATGATTTACAGATCTACGAGCTTACGGAGAGACAGCATAAGAGTTCTGAGTGGAGAACATTATGAACTG 2800
 830 S F L V R A G K P Y D L Q I Y P O E R H S I R V P E S G E H Y E L K 863
 2801 ATCTTTGCAACTACCTTCAAGAAACCTTGGATCACGTTATGCTCTAAAGTGTATTAATTCTGAGCTGAGCTGAGCTGAGCTGAGCTGAGCTG 2900
 863 L L H Y L Q E N L G S R I A A L K V I * 882
 2901 TTAAACCAAATGAGGAGTTTAATCAACAGAAAACAGAAATGATCATCACATTGATACCTGAGGAACTGAGCTGAGCTGAGCTGAGCTGAGCTG 3000
 3001 CCATCACGGCTGAGGAGCTGAGGAGCTGAGGAGCTGAGGAGCTGAGGAGCTGAGGAGCTGAGGAGCTGAGGAGCTGAGGAGCTGAGGAGCTG 3100
 3101 AGAATTACTA[REDACTED] 3127

FIGURE 2

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FIGURE 3
SUBSTITUTE SHEET (RULE 26) RO/AU

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**FIGURE 4**

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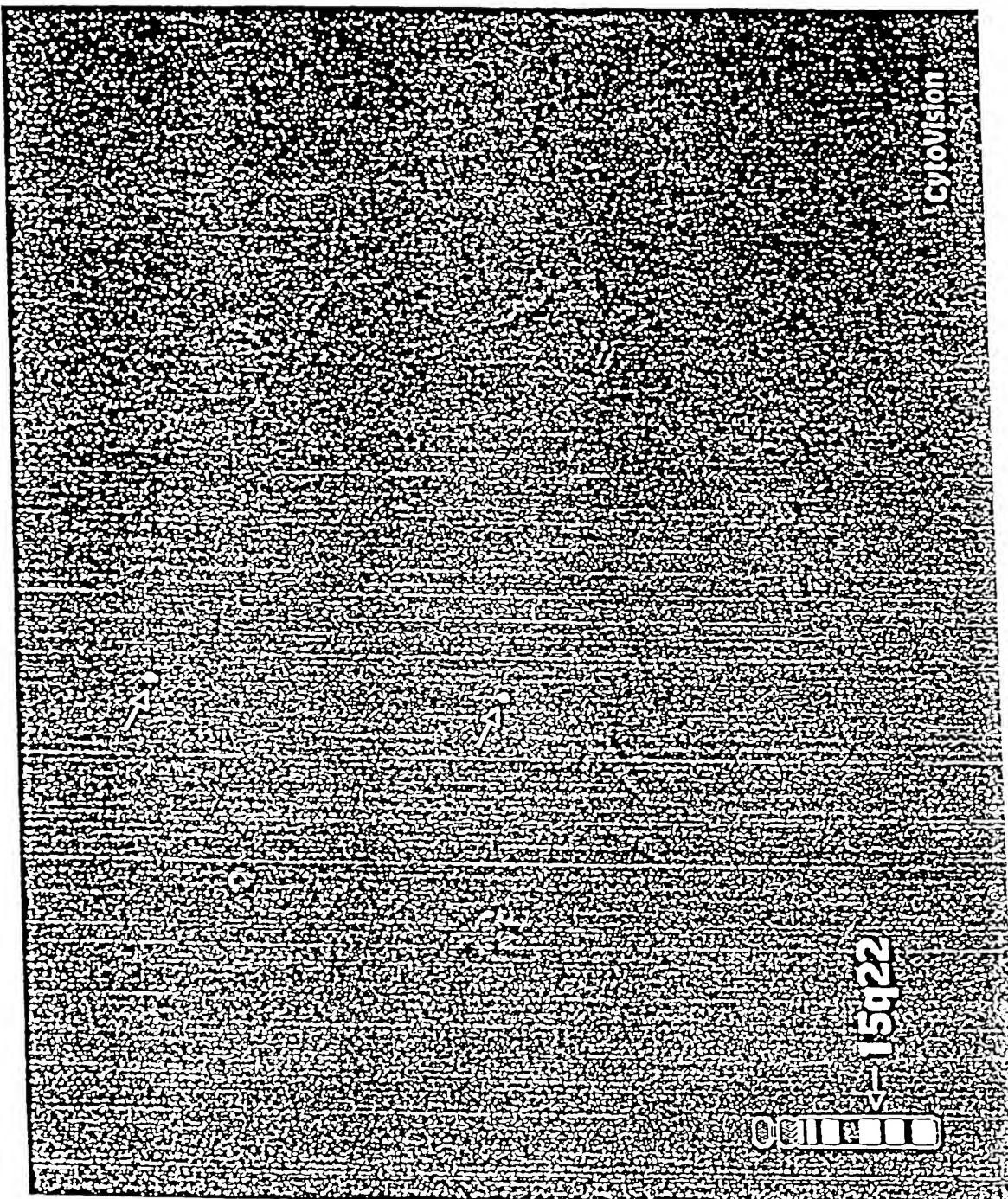


FIGURE 5

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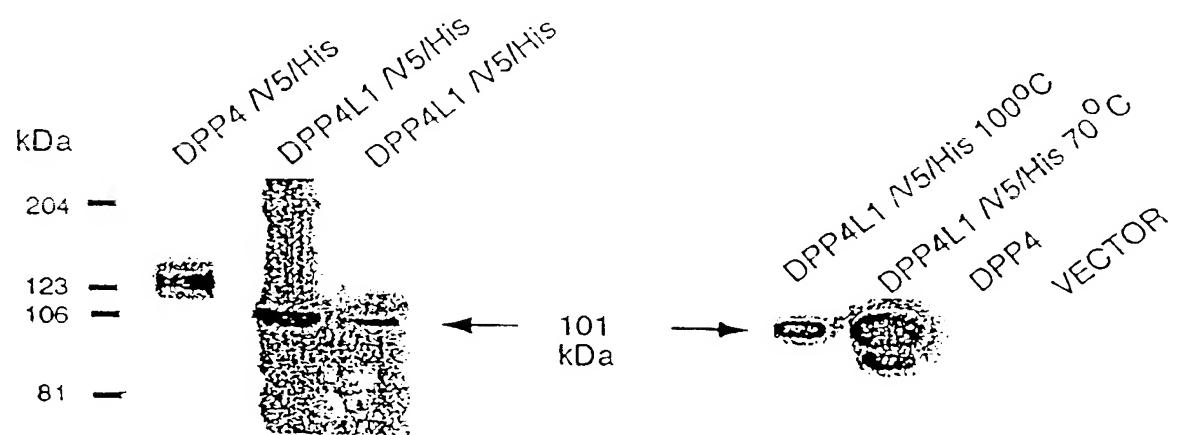


FIGURE 6

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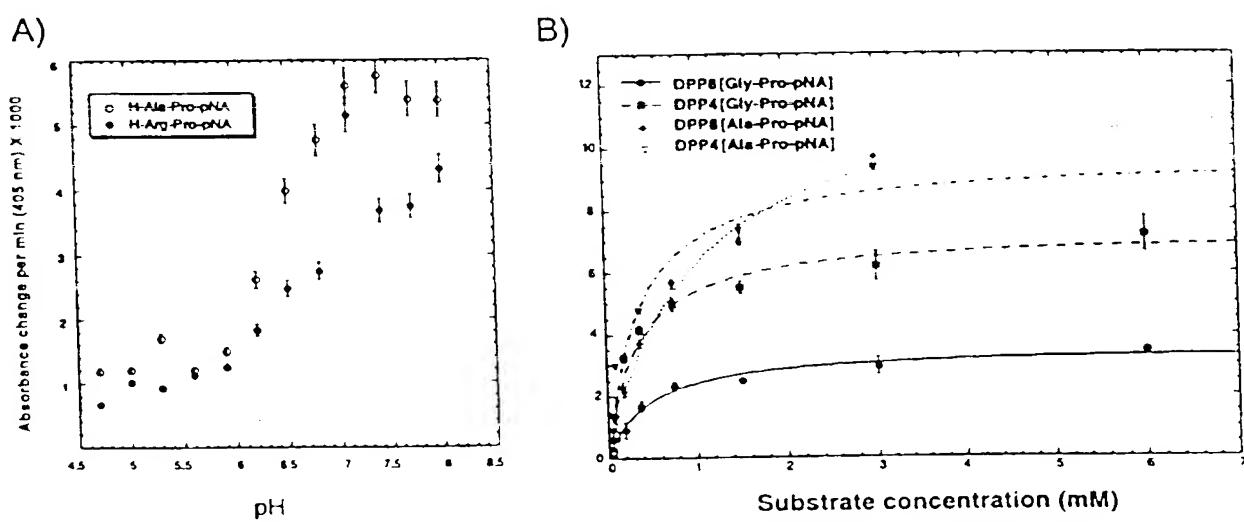


FIGURE 7

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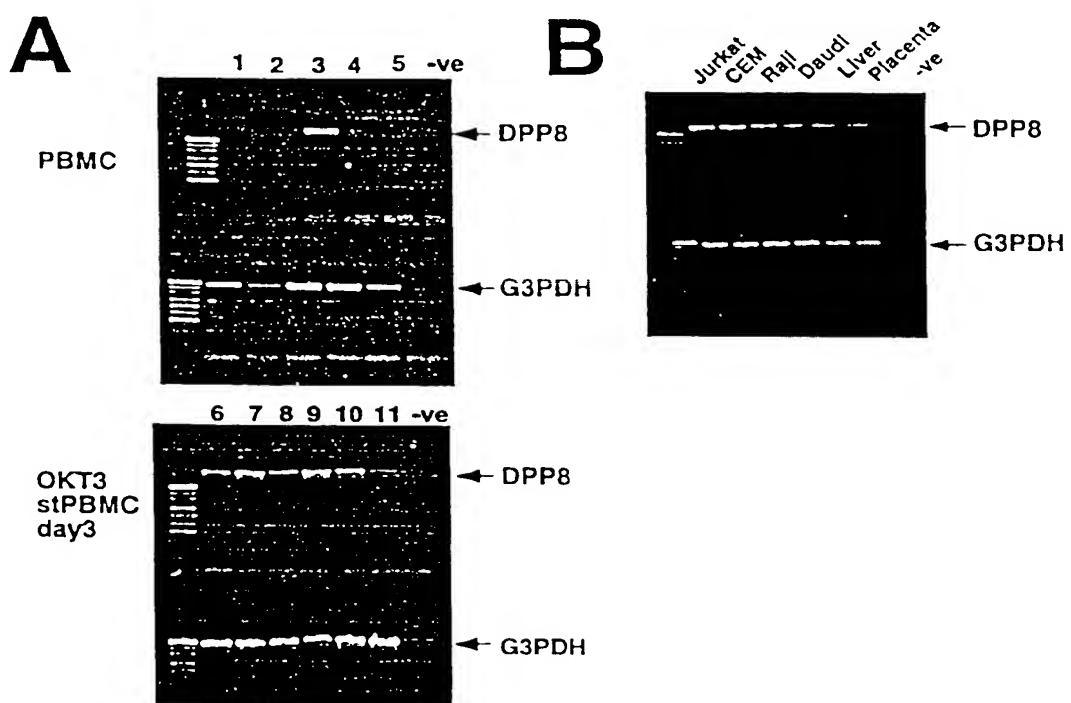


FIGURE 8

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FIGURE 9A

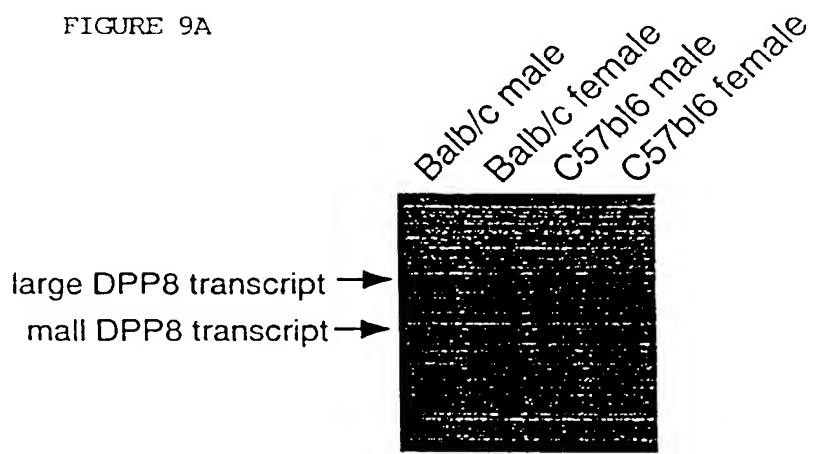


FIGURE 9B

